A Data and Model Visualization System for Android Malware Detection

CPSC 547
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- Rely on machine learning

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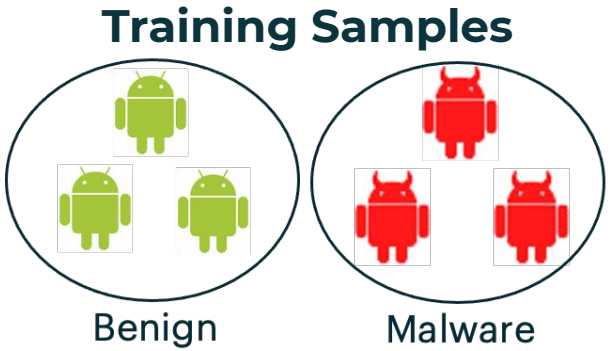

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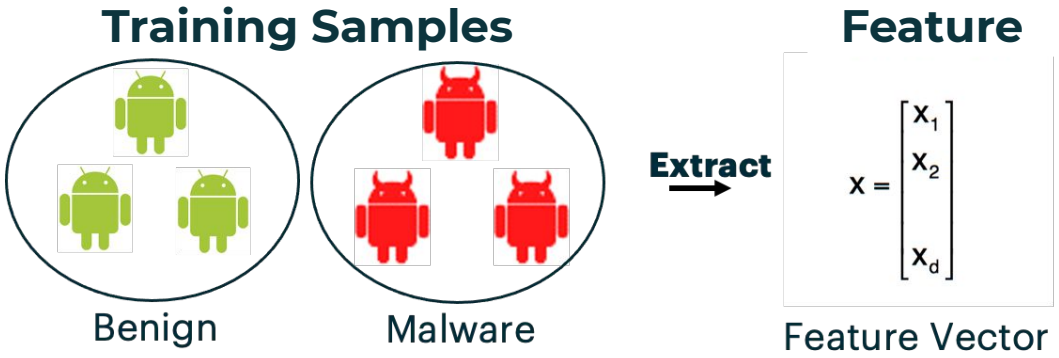

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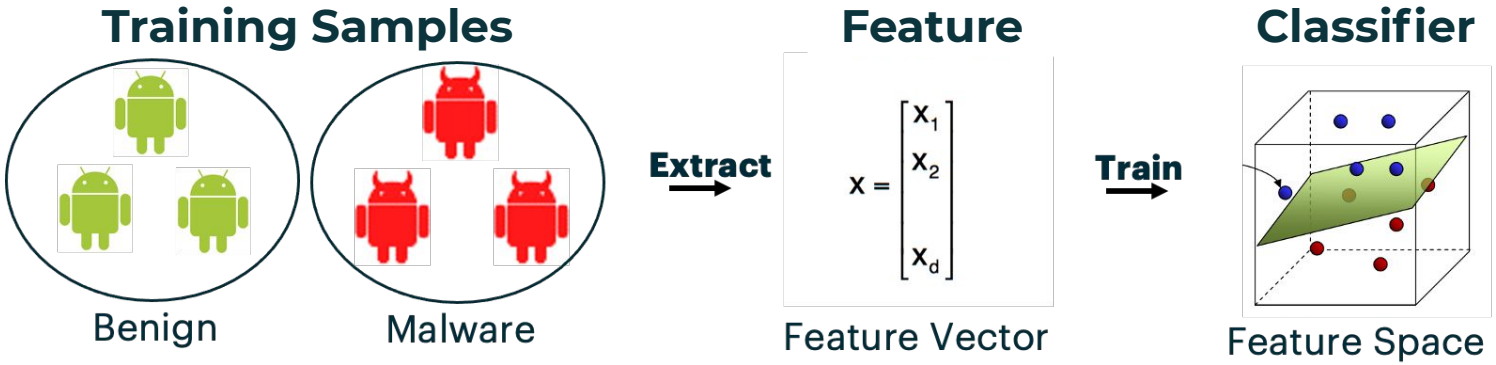

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- Focus on producing models with high accuracy
  - What about model attackability?

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Attackability of the Model is also Important!

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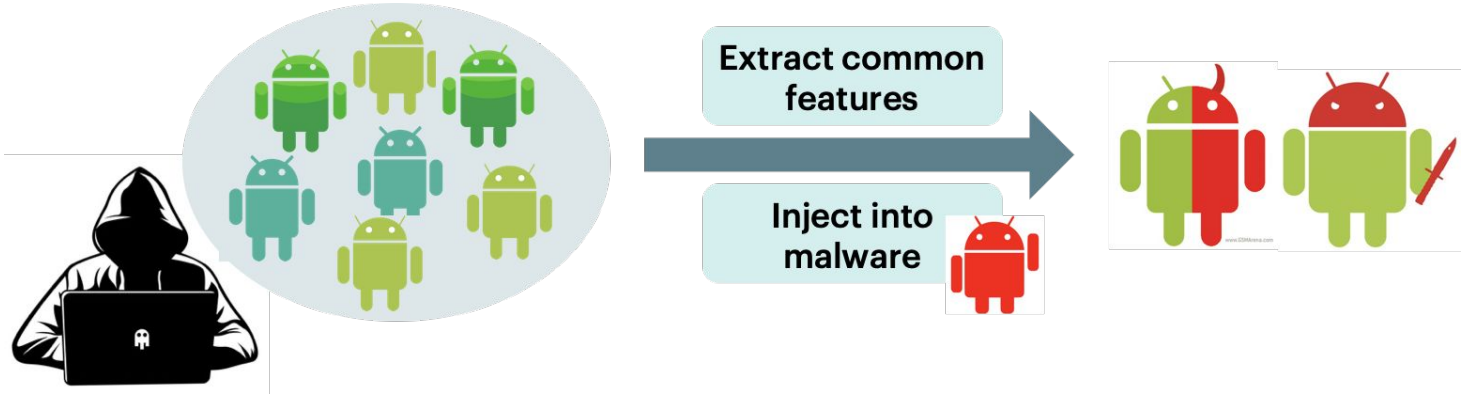
  - What about model attackability?
- Mimicry Attack:
  - Inject features they think represent benign to mislead detection



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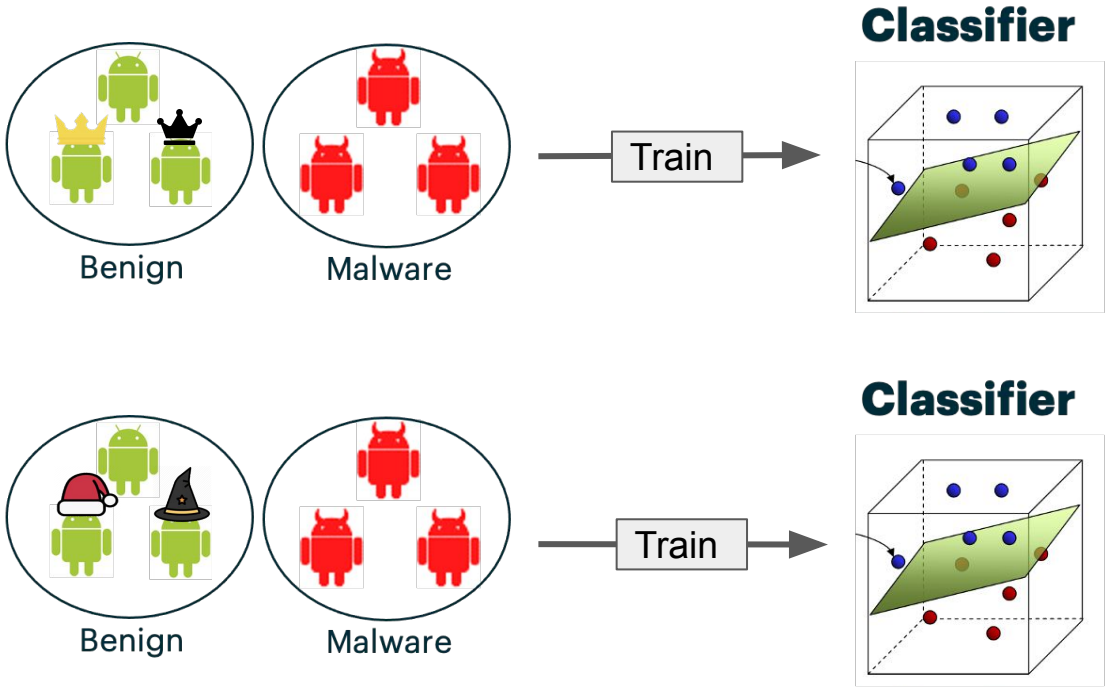

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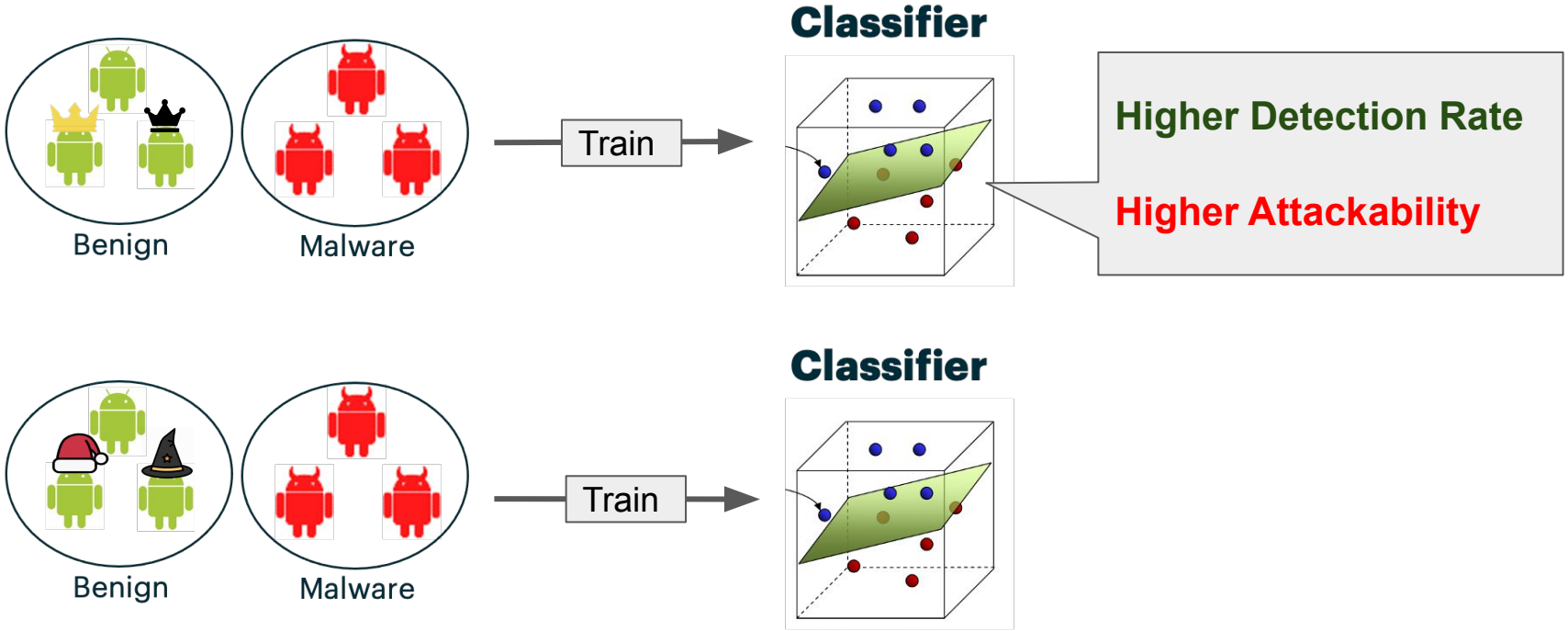

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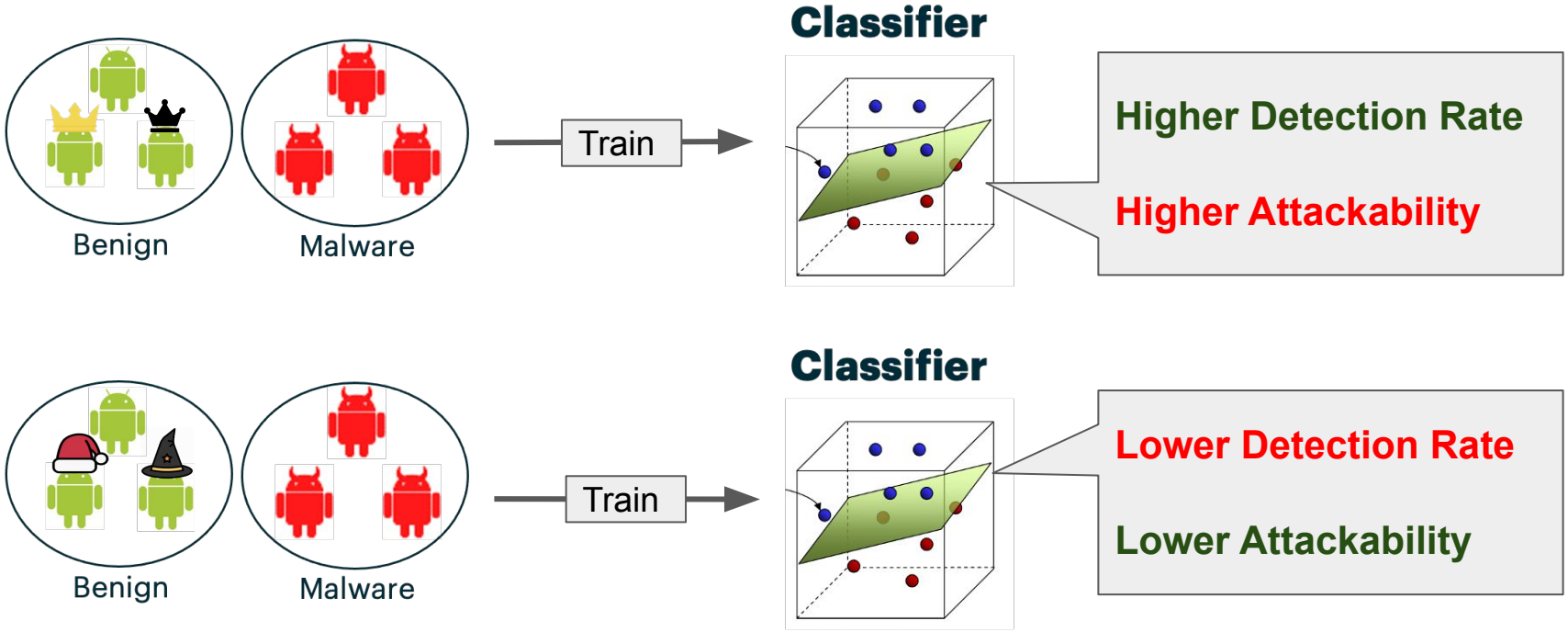

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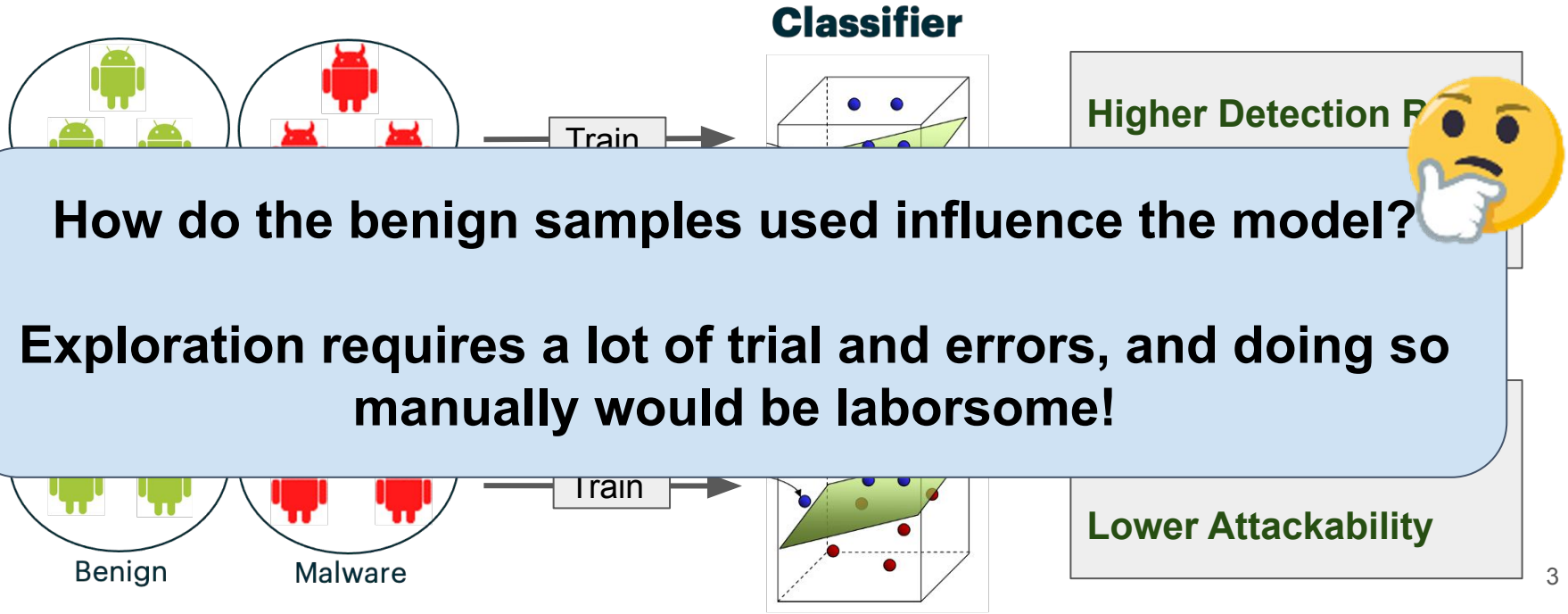

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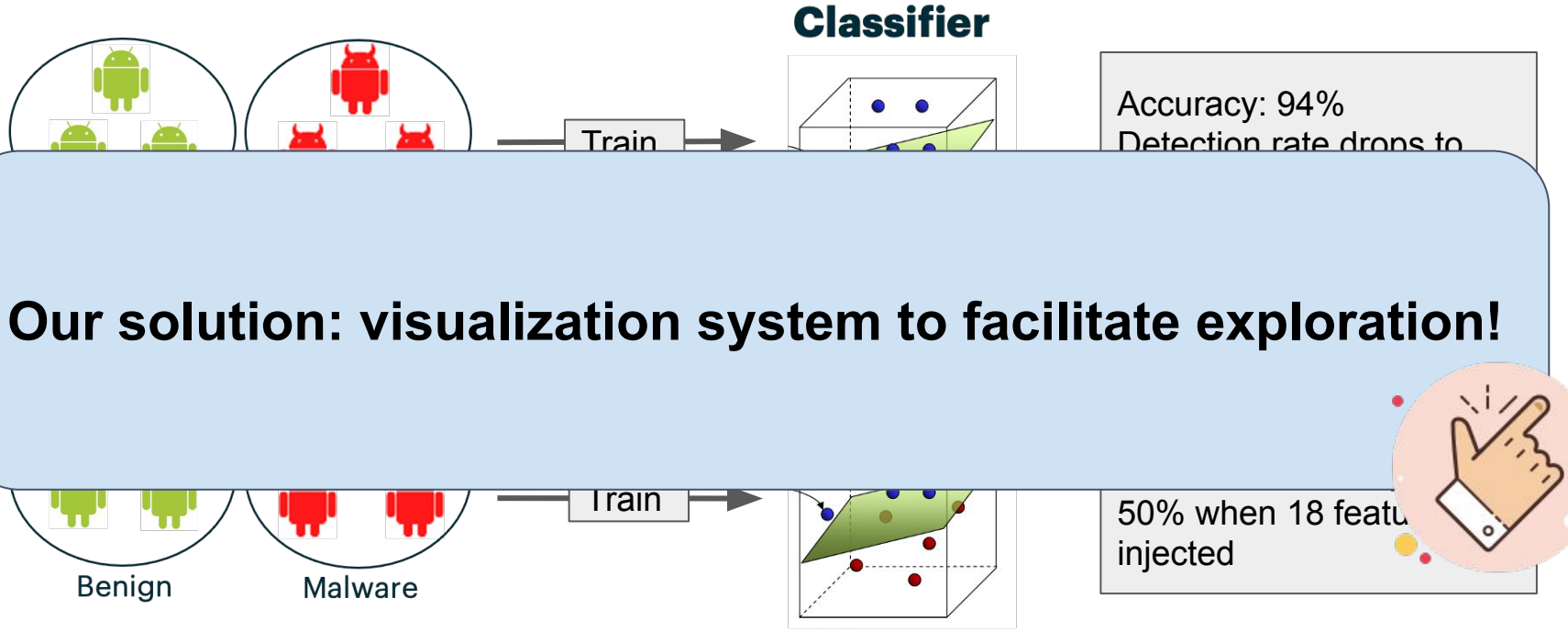
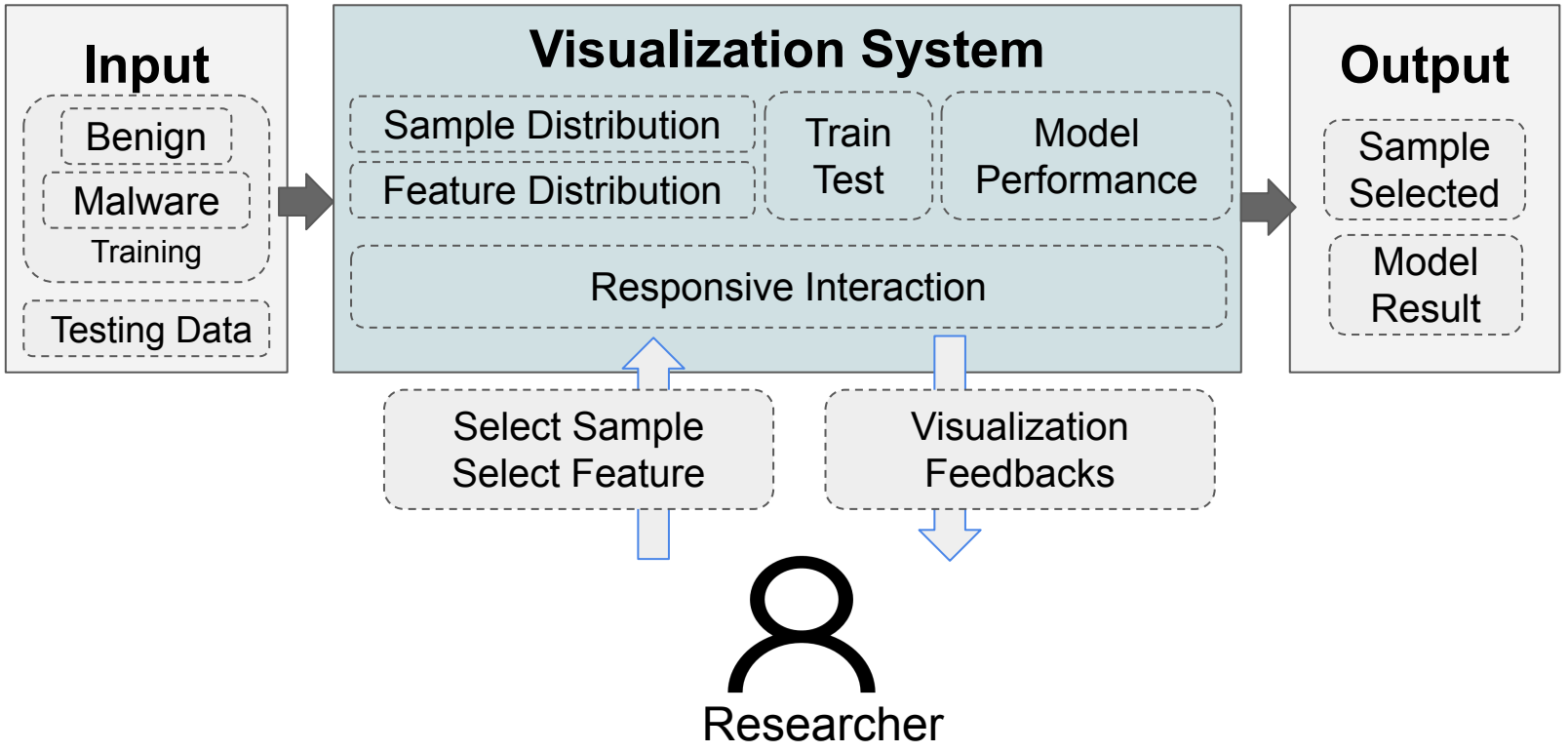


Our solution: visualization system to facilitate exploration!

High Level Overview



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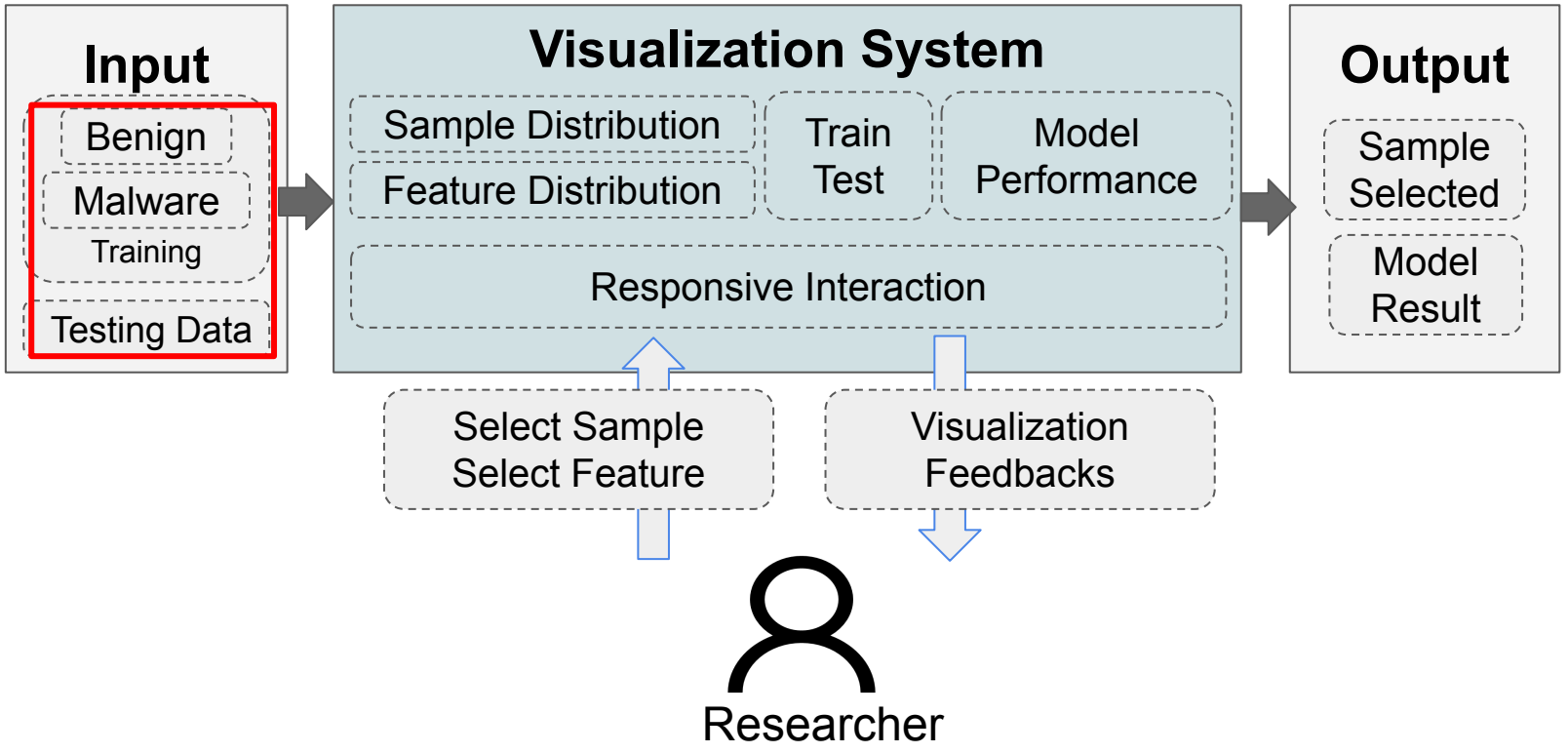

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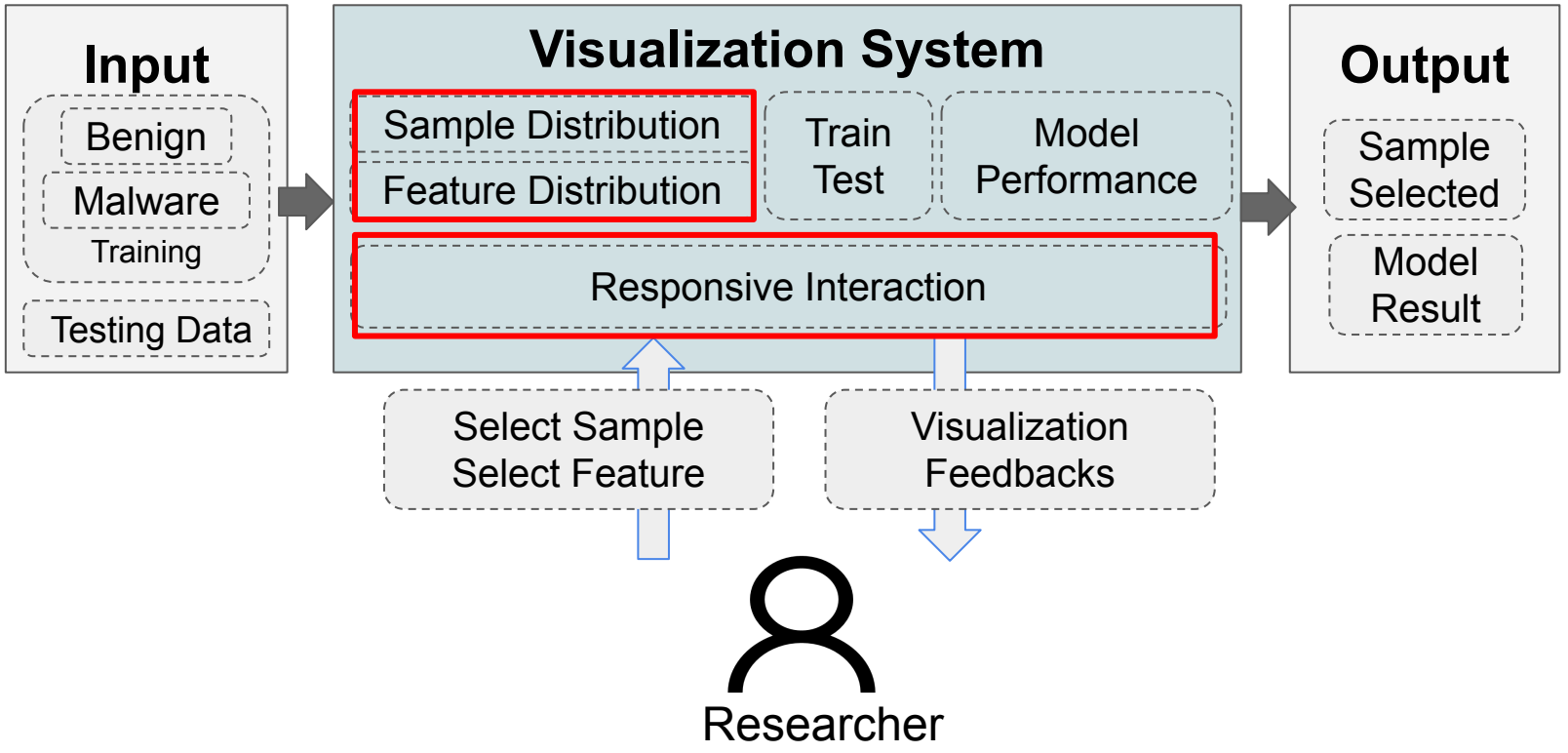

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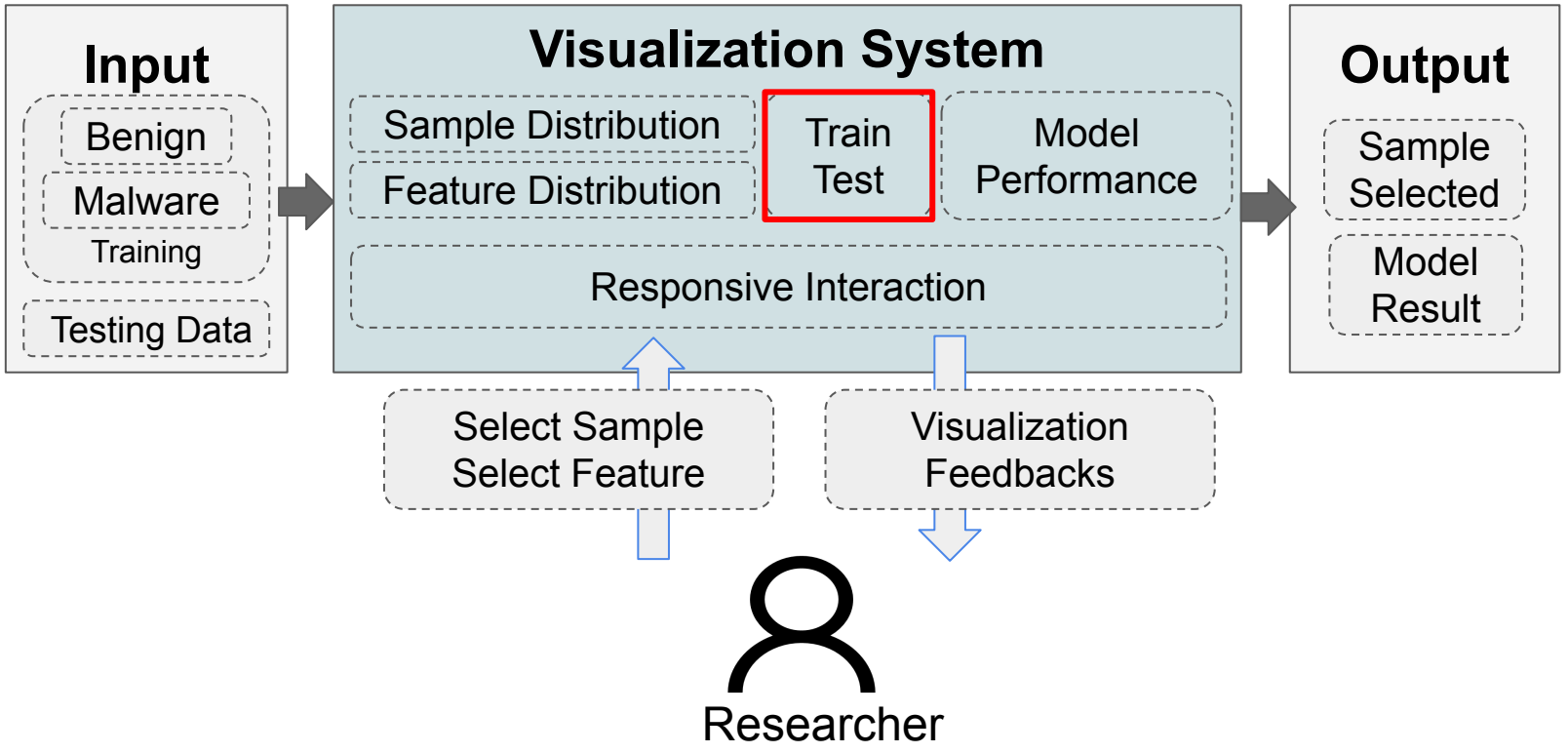

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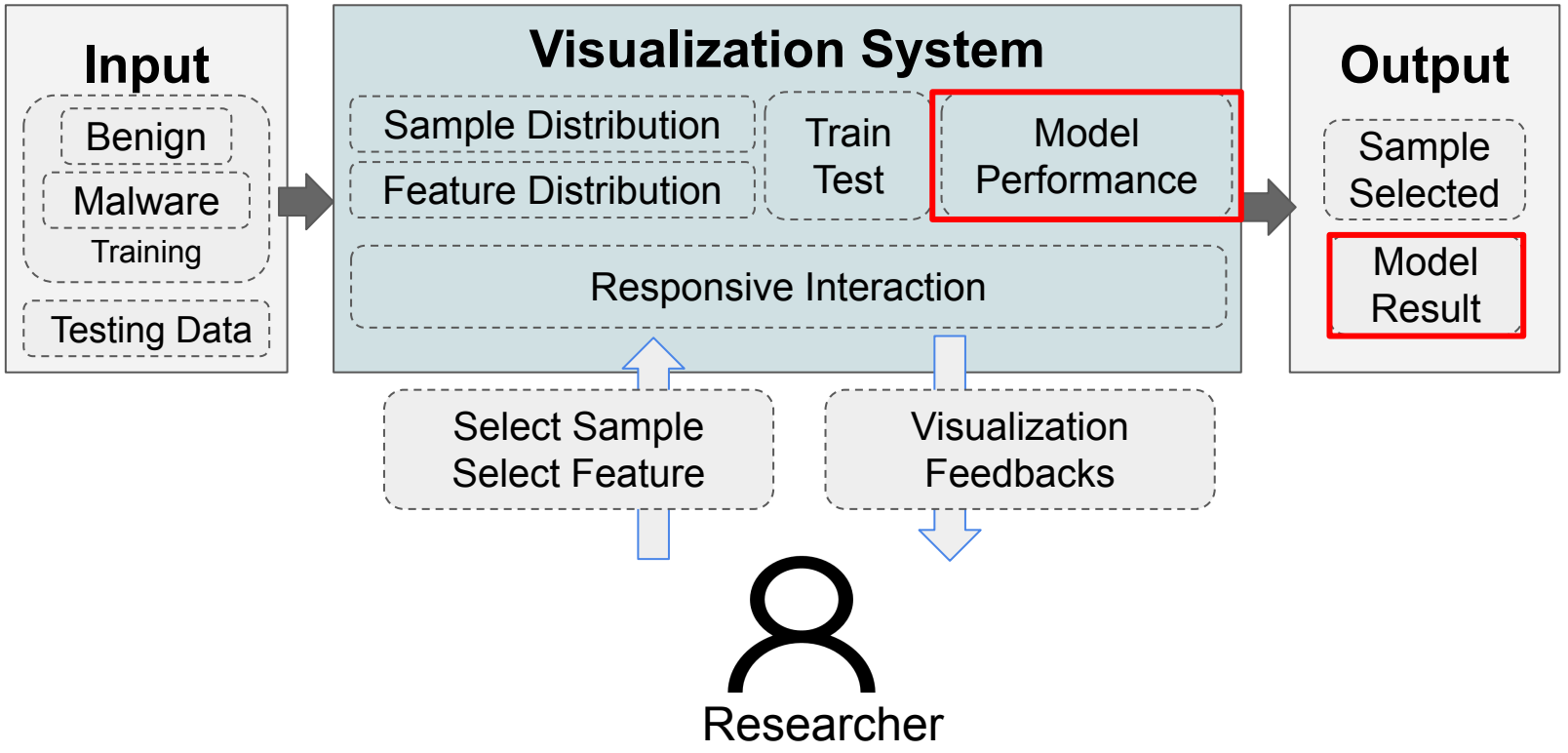

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- Well-known Android malware detection technique [1]
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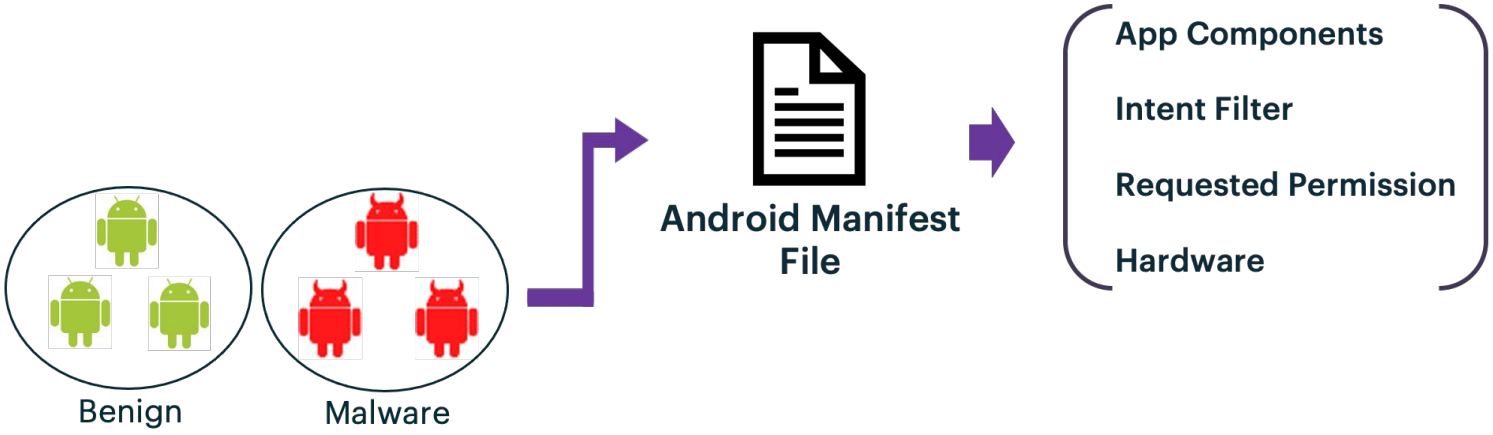

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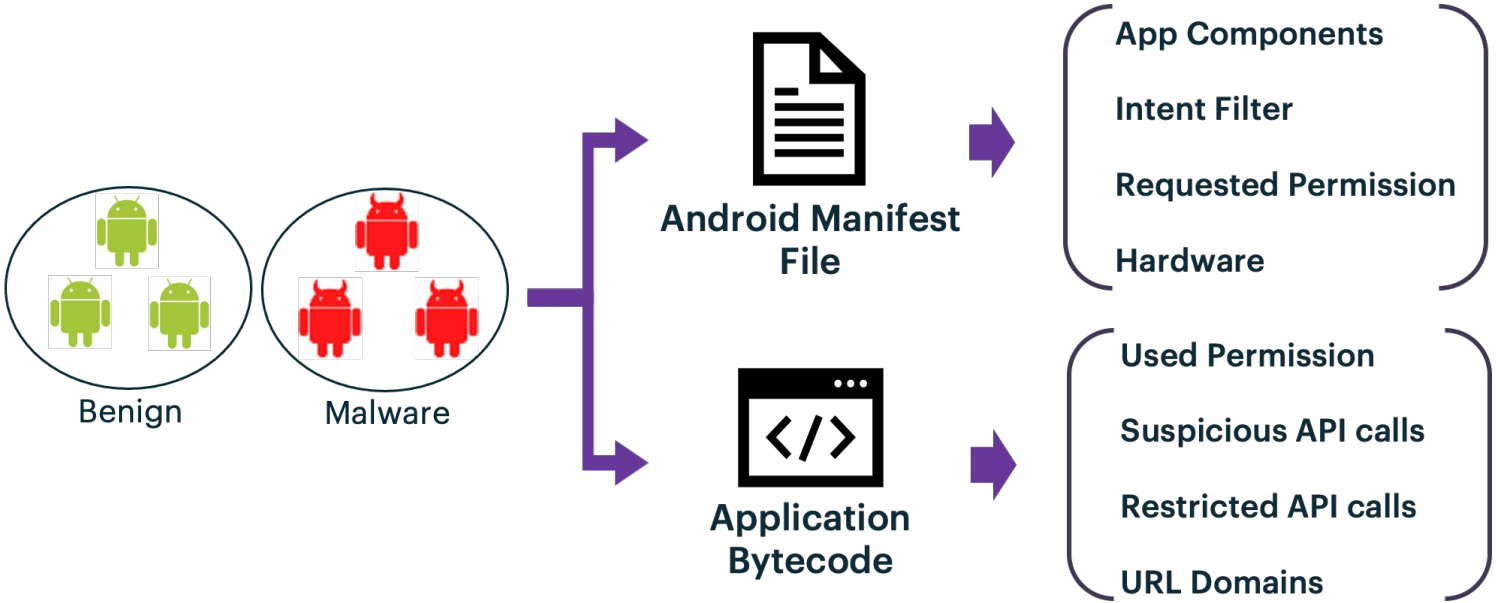
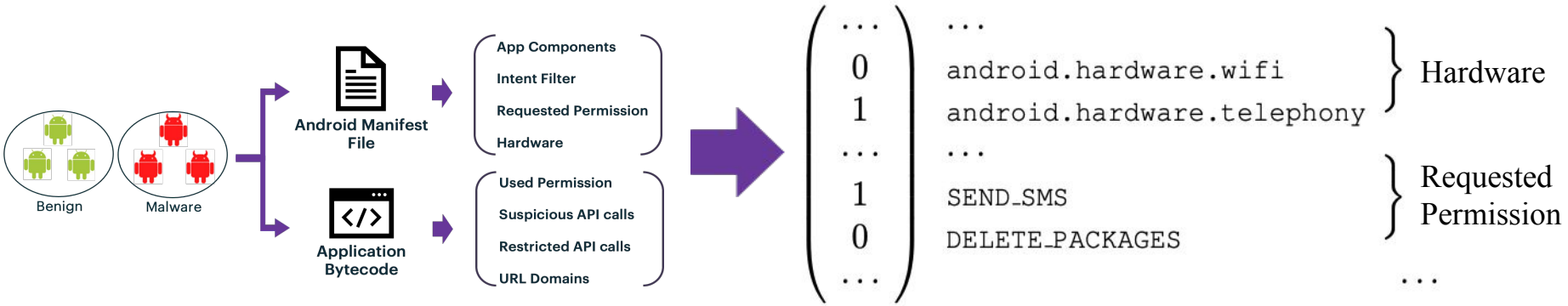
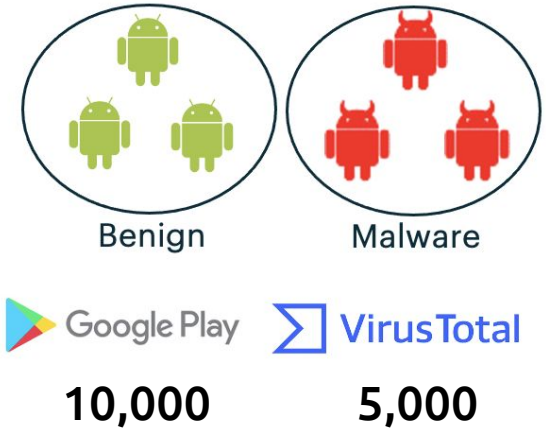


DREBIN: Case Study

- Binary values indicate presence / absence of feature
- Concatenate features from all samples to form feature space
- SVM classifier



Data



Android Sample Data:

- 5,000 malware from VirusTotal
- 10,000 benign from Google Play
- From year 2011 to 2019

Types of Data:

- Android Samples
  - Temporal
  - Drebin Features
- Model
  - Performance
  - Attackability

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- Select particular set of samples and features to train model

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    - Locate misclassified samples
  - Interpret why certain samples being misclassified

Demo

Working scenario of the tool

Limitations

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- Dealing with large feature space
  - Scalability
  - More flexible approaches to select features
  - User has no idea on features for particular samples

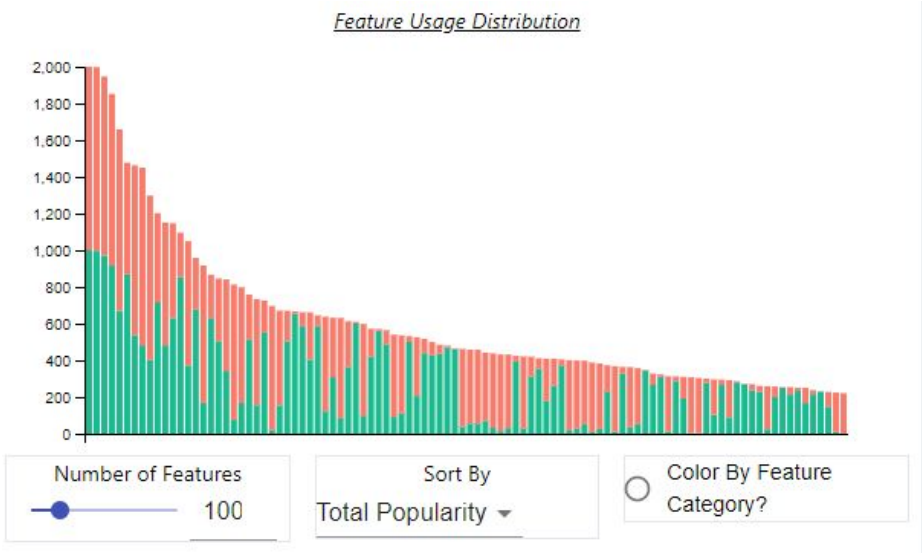
Limitations

- No mechanism for cross experiment comparisons
  - Add juxtaposed view for comparisons
  - Less control over the testing samples

- Few “What-If” functionalities included in the system
  - Allow user to modify / oversample training samples

Lesson Learned

- Prototype, Prototype, Prototype!



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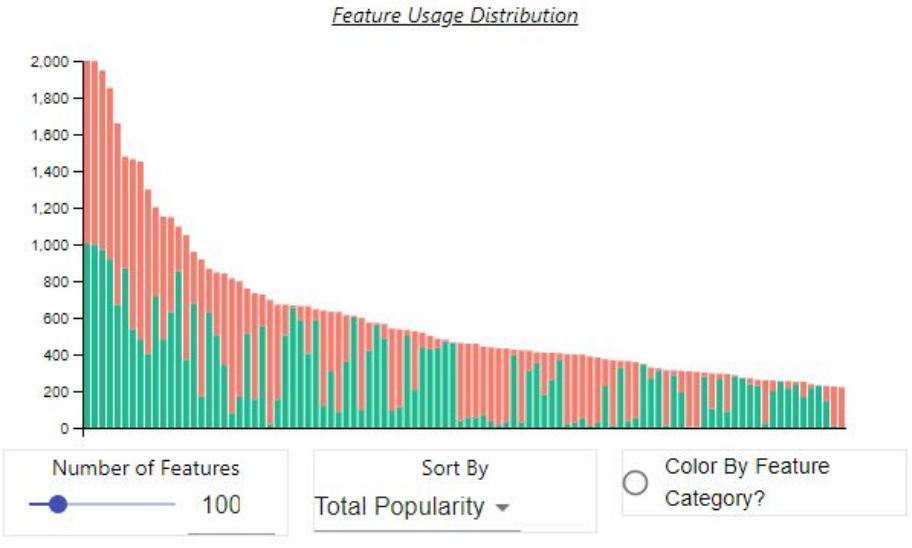

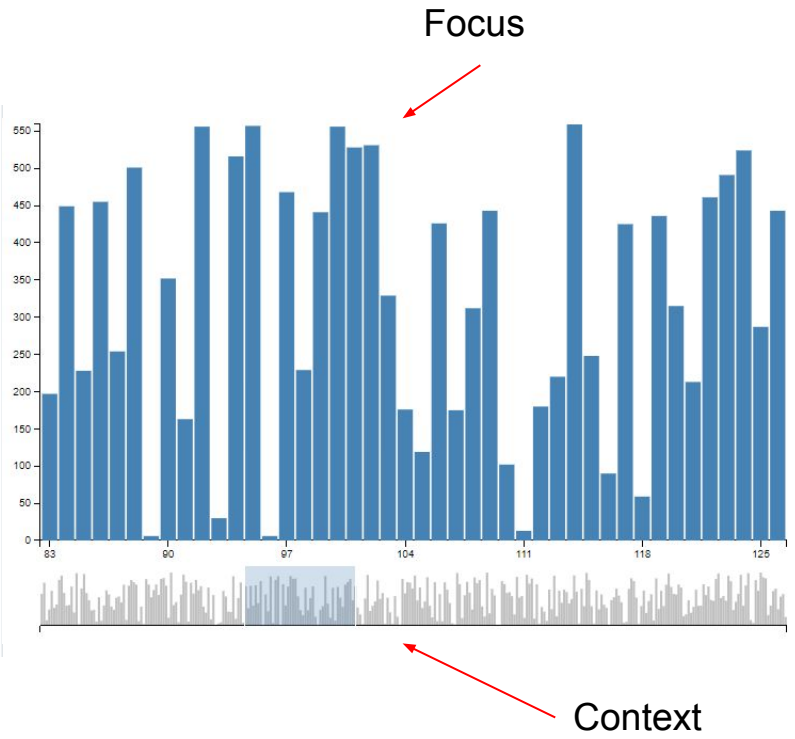

Thank you! Q&A

Intro & Framing

- Research Topic (a tool to facilitate exploring the relationship between training data and resulting model)?
  - How to select the set of benign samples that results in the best performance?
  - OR
  - Current process of performing such exploratory tasks are time consuming?